



## SEQUENCE LISTING

<110> ASHIKARI, TOSHIHIKO  
TANAKA, YOSHIKAZU  
FUJIWARA, HIROYUKI  
NAKAO, MASAHIRO  
FUKUI, YUKO  
SAKAKIBARA, KEIKO  
MIZUTANI, MASAKO  
KUSUMI, TAKAAKI

<120> A GENE ENCODING A PROTEIN HAVING ACYL GROUP TRANSFER ACTIVITY

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<140> 08/894,356  
<141> 1997-08-18

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<151> 1995-02-17

<150> JP 7-196915  
<151> 1995-06-29

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Gln Gly Lys Asn Gly Arg Ser Ile Asp Val Glu Ile Ser Leu Glu	
415 420 425 430	
gca aat gct atg gag agg ttg gag aaa gat aaa gag ttc ctc atg gaa	1404
Ala Asn Ala Met Glu Arg Leu Glu Lys Asp Lys Glu Phe Leu Met Glu	
435 440 445	
act gct taatttgctt agcttggact caactggcta cactttatTTT atgagctgct	1460
Thr Ala	
atgactcaca tgcatgtatg tttatTTTT ttggaggGGGt tctttcTTT tattgtttc	1520
tatgtttttt ctttcttGta cgttatGAAG agaaACCGAG tataaAGGAA taatgtttc	1580
agtttattaaa aaaaaaaaaaaa aaaaa	1605

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<210> 4
<211> 1479
<212> DNA
<213> Perilla ocimoides

<220>
<221> CDS
<222> (3)...(1340)

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    Val Ile Glu Thr Cys Arg Val Gly Pro Pro Pro Asp Ser Val Ala
     1          5           10          15
gag caa tcg gtg ccg ctc aca ttc ttc gac atg acg tgg ctg cat ttt      95
    Glu Gln Ser Val Pro Leu Thr Phe Phe Asp Met Thr Trp Leu His Phe
     20         25           30
cat ccc atg ctt cag ctc ctc tac gaa ttc cct tgt tcc aag caa      143
    His Pro Met Leu Gln Leu Leu Phe Tyr Glu Phe Pro Cys Ser Lys Gln
     35         40           45
cat ttt tca gaa tcc atc gtt cca aaa ctc aaa caa tct ctc tct aaa      191
    His Phe Ser Glu Ser Ile Val Pro Lys Leu Lys Gln Ser Leu Ser Lys
     50         55           60
act ctc ata cac ttc ttc cct ctc tca tgc aat tta atc tac cct tca      239
    Thr Leu Ile His Phe Phe Pro Leu Ser Cys Asn Leu Ile Tyr Pro Ser
     65         70           75
tcc ccg gag aaa atg ccg gag ttt cgg tat cta tcc ggg gac tcg gtt      287
    Ser Pro Glu Lys Met Pro Glu Phe Arg Tyr Leu Ser Gly Asp Ser Val
     80         85           90           95
tct ttc acc atc gca gaa tct agc gac gac ttc gat gat ctc gtc gga      335
    Ser Phe Thr Ile Ala Glu Ser Ser Asp Asp Phe Asp Asp Leu Val Gly
     100        105          110
aat cgt cca gaa tct ccc gtt agg ctc tac aac ttt gtc cct aaa ttg      383
    Asn Arg Pro Glu Ser Pro Val Arg Leu Tyr Asn Phe Val Pro Lys Leu
     115        120          125
ccg ccc att gtc gaa gaa tcc gat aga aaa ctc ttc caa gtt ttc gcc      431
    Pro Pro Ile Val Glu Ser Asp Arg Lys Leu Phe Gln Val Phe Ala
     130        135          140
gtg cag gtg act ctt ttc cca ggc cga ggc gtc ggt att gga ata gca      479
    Val Gln Val Thr Leu Phe Pro Gly Arg Gly Val Gly Ile Gly Ile Ala
     145        150          155
acg cat cac acc gtt agc gac gcc ccg tcg ttt ctc gcg ttt ata acg      527
    Thr His His Thr Val Ser Asp Ala Pro Ser Phe Leu Ala Phe Ile Thr
     160        165          170          175
gct tgg tct tca atg agc aaa cac att gaa aat gaa gat gaa gat gaa      575
    Ala Trp Ser Ser Met Ser Lys His Ile Glu Asn Glu Asp Glu Asp Glu
     180        185          190

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gaa ttt aaa tct ttg cca gtt ttc gat aga tcc gtc ata aaa tat ccg Glu Phe Lys Ser Leu Pro Val Phe Asp Arg Ser Val Ile Lys Tyr Pro 195 200 205	623
acg aaa ttt gac tcc att tat tgg aga aac gcg cta aaa ttt cct ttg Thr Lys Phe Asp Ser Ile Tyr Trp Arg Asn Ala Leu Lys Phe Pro Leu 210 215 220	671
caa tct cgt cat ccc tca tta ccg acg gac cgc att cga acc acg ttc Gln Ser Arg His Pro Ser Leu Pro Thr Asp Arg Ile Arg Thr Thr Phe 225 230 235	719
gtt ttc acc caa tcc aaa att aag aaa ttg aag ggt tgg att cag tcc Val Phe Thr Gln Ser Lys Ile Lys Lys Leu Lys Gly Trp Ile Gln Ser 240 245 250 255	767
aga gtt cca agt tta gtc cat ctc tca tct ttt gta gcg att gca gct Arg Val Pro Ser Leu Val His Leu Ser Ser Phe Val Ala Ile Ala Ala 260 265 270	815
tat atg tgg gct ggc ata acg aaa tca ttc aca gca gat gaa gac caa Tyr Met Trp Ala Gly Ile Thr Lys Ser Phe Thr Ala Asp Glu Asp Gln 275 280 285	863
gac aac gag gat gca ttt ttc ttg att ccg gtc gat cta agg cca cga Asp Asn Glu Asp Ala Phe Phe Leu Ile Pro Val Asp Leu Arg Pro Arg 290 295 300	911
tta gat ccg ccg gtt cct gaa aat tac ttc ggg aac tgc tta tcg tac Leu Asp Pro Pro Val Pro Glu Asn Tyr Phe Gly Asn Cys Leu Ser Tyr 305 310 315	959
gcg ctg ccg aga atg cgg cgg cga gag ctg gtg gga gag aaa ggg gtg Ala Leu Pro Arg Met Arg Arg Glu Leu Val Gly Glu Lys Gly Val 320 325 330 335	1007
ttt ctg gca gct gag gta atc gcg gcg gag ata aaa agg atc aac Phe Leu Ala Ala Glu Val Ile Ala Ala Glu Ile Lys Lys Arg Ile Asn 340 345 350	1055
gac aag aga ata tta gaa acg gtg gag aaa tgg tcg ccg gag att cgt Asp Lys Arg Ile Leu Glu Thr Val Glu Lys Trp Ser Pro Glu Ile Arg 355 360 365	1103
aaa gcg ttg cag aaa tca tat ttt tcg gtg gca gga tcg agc aag cta Lys Ala Leu Gln Lys Ser Tyr Phe Ser Val Ala Gly Ser Ser Lys Leu 370 375 380	1151
gat ctt tac ggt gca gat ttt gga tgg ggg aag gcg aga aag caa gaa Asp Leu Tyr Gly Ala Asp Phe Gly Trp Gly Lys Ala Arg Lys Gln Glu 385 390 395	1199
ata ttg tcg att gat ggg gag aaa tat gca atg acg ctt tgt aaa gcc Ile Leu Ser Ile Asp Gly Glu Lys Tyr Ala Met Thr Leu Cys Lys Ala 400 405 410 415	1247

agg gat ttc gaa gga gga ttg gag gtt tgc ttg tct ttg cct aag gac	1295
Arg Asp Phe Glu Gly Gly Leu Glu Val Cys Leu Ser Leu Pro Lys Asp	
420	425
430	
aaa atg gat gct ttt gct tat ttt tca ctg gga att aat ggt	1340
Lys Met Asp Ala Phe Ala Ala Tyr Phe Ser Leu Gly Ile Asn Gly	
435	440
445	
taataaaatgt atgtaattaa actaatatta ttatgtaaca attaattaag tgttgagtaa	1400
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tgaaaaaaaaaaaaaaa	1479

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<213> Senecio cruentus	
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<222> (3) .. (1364)	
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Asn Ile Leu Glu His Ala Arg Ile Ser Ala Pro Ser Gly Thr Ile	
1	5
10	15
ggc cat cgc tcg tta tct ctt act ttc ttc gac att act tgg cta ctc	95
Gly His Arg Ser Leu Ser Leu Thr Phe Phe Asp Ile Thr Trp Leu Leu	
20	25
30	
ttc cct ccg gtc cac cat ctt ttc tat gac ttt cca cat tct aaa	143
Phe Pro Pro Val His His Leu Phe Phe Tyr Asp Phe Pro His Ser Lys	
35	40
45	
tcc cat ttc atg gac act att gtt ccc agg cta aaa caa tct tta tcg	191
Ser His Phe Met Asp Thr Ile Val Pro Arg Leu Lys Gln Ser Leu Ser	
50	55
60	
gtc act ctt caa cat ttt ttc ccg ttt gct agt aat ttg att gta ttt	239
Val Thr Leu Gln His Phe Phe Pro Phe Ala Ser Asn Leu Ile Val Phe	
65	70
75	
cct aac act gat ggt tcg ggt ttt aat aaa aaa cca gaa ata aaa cac	287
Pro Asn Thr Asp Gly Ser Gly Phe Asn Lys Lys Pro Glu Ile Lys His	
80	85
90	95
gtt gaa ggt gat tct gtg gtt act ttt gca gaa tgt tgt ctt gac	335
Val Glu Gly Asp Ser Val Val Val Thr Phe Ala Glu Cys Cys Leu Asp	
100	105
110	
ttt aat aat ttg aca gga aat cat cct cga aaa tgt gaa aac ttt tat	383
Phe Asn Asn Leu Thr Gly Asn His Pro Arg Lys Cys Glu Asn Phe Tyr	
115	120
125	

cca ctt gta cct tca ttg gga aat gca atc aaa tta tgt gat tgc gtc	431
Pro Leu Val Pro Ser Leu Gly Asn Ala Ile Lys Leu Cys Asp Cys Val	
130 135 140	
acg gtc cca ctt ttt tca ctt caa gtg acg ttt ttt ccg ggc tcg ggt	479
Thr Val Pro Leu Phe Ser Leu Gln Val Thr Phe Phe Pro Gly Ser Gly	
145 150 155	
ata tca cta gga atg acg aat cat cat agc ctt ggt gac gct agc acg	527
Ile Ser Leu Gly Met Thr Asn His His Ser Leu Gly Asp Ala Ser Thr	
160 165 170 175	
cggttc aac ttt ttg aaa ggg tgg act tcg att att caa tct ggt gta	575
Arg Phe Asn Phe Leu Lys Gly Trp Thr Ser Ile Ile Gln Ser Gly Val	
180 185 190	
gat cgg tct ttt tta acg aaa gga tct cca ccg gtt ttt gat aga ttg	623
Asp Arg Ser Phe Leu Thr Lys Gly Ser Pro Pro Val Phe Asp Arg Leu	
195 200 205	
att aac atc cca cat tta gat gaa aat aag ttg aga cat aca agg ctc	671
Ile Asn Ile Pro His Leu Asp Glu Asn Lys Leu Arg His Thr Arg Leu	
210 215 220	
gaa agt ttt tat aaa cct tcg agc ctt gtt ggt ccc act gat aaa gtt	719
Glu Ser Phe Tyr Lys Pro Ser Ser Leu Val Gly Pro Thr Asp Lys Val	
225 230 235	
cggtca acg ttt gtg ttg acc cga act aat atc aat cta cta aag aaa	767
Arg Ser Thr Phe Val Leu Thr Arg Thr Asn Ile Asn Leu Leu Lys Lys	
240 245 250 255	
aag gtc tta acc caa gtg cca aac ttg gag tac atg tca tct ttt acg	815
Lys Val Leu Thr Gln Val Pro Asn Leu Glu Tyr Met Ser Ser Phe Thr	
260 265 270	
gta act tgt ggt tat ata tgg agt tgc ata gcg aaa tca ctc gta aaa	863
Val Thr Cys Gly Tyr Ile Trp Ser Cys Ile Ala Lys Ser Leu Val Lys	
275 280 285	
ata gga gaa aga aag ggc gaa gac gag tta gaa cag ttc ata atc acc	911
Ile Gly Glu Arg Lys Gly Glu Asp Glu Leu Glu Gln Phe Ile Ile Thr	
290 295 300	
att gat tgt cga tct cgt ctt gat cca cca att ccc aca gcc tac ttt	959
Ile Asp Cys Arg Ser Arg Leu Asp Pro Pro Ile Pro Thr Ala Tyr Phe	
305 310 315	
ggtaac tgt ggt gca cca tgt gtc ccg acc tta aaa aat gtc gtt ttg	1007
Gly Asn Cys Gly Ala Pro Cys Val Pro Thr Leu Lys Asn Val Val Leu	
320 325 330 335	
act acg gaa aat ggg tat gca ctt ggt gct aaa gta att gga gag tct	1055
Thr Thr Glu Asn Gly Tyr Ala Leu Gly Ala Lys Val Ile Gly Glu Ser	
340 345 350	

ata tgc aaa atg ata tat aat aag gac gga atc ttg aaa gat gcc gcg Ile Cys Lys Met Ile Tyr Asn Lys Asp Gly Ile Leu Lys Asp Ala Ala 355 360 365	1103
aga tgg cat gaa cct ttc atg atc ccg gct agg aag att ggt gtt gct Arg Trp His Glu Pro Phe Met Ile Pro Ala Arg Lys Ile Gly Val Ala 370 375 380	1151
ggt aca cct aag ctc aac ttg tac gac ttt gat ttt ggg tgg ggg aag Gly Thr Pro Lys Leu Asn Leu Tyr Asp Phe Asp Phe Gly Trp Gly Lys 385 390 395	1199
cgc ata aag tat gag act gtt tca ata gac tat aat acg tcg att tct Arg Ile Lys Tyr Glu Thr Val Ser Ile Asp Tyr Asn Thr Ser Ile Ser 400 405 410 415	1247
ata aat gca agc aaa aca tca gca caa gat ctt gaa att gga ttg agt Ile Asn Ala Ser Lys Thr Ser Ala Gln Asp Leu Glu Ile Gly Leu Ser 420 425 430	1295
cta ccg agt atg caa atg gag gcg ttt tct agc atc ttt gat gaa gga Leu Pro Ser Met Gln Met Glu Ala Phe Ser Ser Ile Phe Asp Glu Gly 435 440 445	1343
tta gag agt caa gtt tca ttg tagatcatcg tccccctttt gtgtgcata Leu Glu Ser Gln Val Ser Leu 450	1394
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<212> DNA  
<213> Lavandula angustifolia

<220>  
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<222> (3)...(1352)

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tg acc acc ctc ctc gaa tcc tcc cga gtg gcg ccg cct cca ggc acg Thr Thr Leu Leu Glu Ser Ser Arg Val Ala Pro Pro Pro Gly Thr 1 5 10 15	47
gtg gct gag cag tca ctc ccg ctc acc ttc ttc gac atg acg tgg ctg Val Ala Glu Gln Ser Leu Pro Leu Thr Phe Phe Asp Met Thr Trp Leu 20 25 30	95
cat ttc cac ccc atg ctt cag ctc ttc tac gaa ctc ccc tgt tcc His Phe His Pro Met Leu Gln Leu Leu Phe Tyr Glu Leu Pro Cys Ser 35 40 45	143

aaa ccc gcc ttc ctc gaa acc gtc gtt ccg aaa ctc aaa caa tcc tta	191		
Lys Pro Ala Phe Leu Glu Thr Val Val Pro Lys Leu Lys Gln Ser Leu			
50	55	60	
tct cta acc ctc aaa cac ttc ttc ccc ctt tca tgc aat cta atc tac	239		
Ser Leu Thr Leu Lys His Phe Phe Pro Leu Ser Cys Asn Leu Ile Tyr			
65	70	75	
cct cta tcg ccg gag aaa atg ccg gag ttc cgg tat cag aac ggt gac	287		
Pro Leu Ser Pro Glu Lys Met Pro Glu Phe Arg Tyr Gln Asn Gly Asp			
80	85	90	95
tcg gtt tct ttc acg att atg gag tct gtc gga gat cat ccg cat tcc	335		
Ser Val Ser Phe Thr Ile Met Glu Ser Val Gly Asp His Pro His Ser			
100	105	110	
gct cat aaa tac tac tgc ttt gcc cct agc gac gat tat gaa gat ctc	383		
Ala His Lys Tyr Tyr Cys Phe Ala Pro Ser Asp Asp Tyr Glu Asp Leu			
115	120	125	
cag ctg ccg ccg ata gtc gag gaa tct gat cgg aaa ttg tttcaa gtt	431		
Gln Leu Pro Pro Ile Val Glu Ser Asp Arg Lys Leu Phe Gln Val			
130	135	140	
tta gcc gtg caa gtg act ctg ttt ccc ggt cgc ggg gtg tgc atc gga	479		
Leu Ala Val Gln Val Thr Leu Phe Pro Gly Arg Gly Val Cys Ile Gly			
145	150	155	
ata acg acg cac cac acc gtt agc gat gct cca tcg ttt gta ggg ttt	527		
Ile Thr Thr His His Thr Val Ser Asp Ala Pro Ser Phe Val Gly Phe			
160	165	170	175
atg aag agt tgg gct tcc atc act aaa ttc gga gga gat gat gaa ttc	575		
Met Lys Ser Trp Ala Ser Ile Thr Lys Phe Gly Gly Asp Asp Glu Phe			
180	185	190	
ttg gac gga aaa ggt gaa tgt ttg ccg gtt ttc gac cga tcg ctc gtg	623		
Leu Asp Gly Lys Gly Glu Cys Leu Pro Val Phe Asp Arg Ser Leu Val			
195	200	205	
aat tat ccg cct aaa ttg gac aca tat tta tgg aac aac gcg cag aaa	671		
Asn Tyr Pro Pro Lys Leu Asp Thr Tyr Leu Trp Asn Asn Ala Gln Lys			
210	215	220	
cgt ccg ttg gaa tcg cag cat cca tct tta ccg acg gat cgg att cga	719		
Arg Pro Leu Glu Ser Gln His Pro Ser Leu Pro Thr Asp Arg Ile Arg			
225	230	235	
gct acc tac ctt ttc acc caa tct gaa att aag aaa ttg aag ggt ttg	767		
Ala Thr Tyr Leu Phe Thr Gln Ser Glu Ile Lys Lys Leu Lys Gly Leu			
240	245	250	255
att cag aga aaa gcc cca aat gta gtt aat ctc tct tcc ttc gtc gcg	815		
Ile Gln Arg Lys Ala Pro Asn Val Val Asn Leu Ser Ser Phe Val Ala			
260	265	270	

atc gca gct tat atc tgg acc ggc atc gcc aaa tcg gtc gga gat tac Ile Ala Ala Tyr Ile Trp Thr Gly Ile Ala Lys Ser Val Gly Asp Tyr 275 280 285	863
aaa gac gtg gat gac gac aaa cgc gct ttc ttt tta att ccg atc gat Lys Asp Val Asp Asp Lys Arg Ala Phe Phe Leu Ile Pro Ile Asp 290 295 300	911
tta agg ccg cgt ttg gat ccg ccg gct ccg ggg aac tac ttc gga aac Leu Arg Pro Arg Leu Asp Pro Pro Ala Pro Gly Asn Tyr Phe Gly Asn 305 310 315	959
tgt cta tcg ttt gcg atg gcg aag atc ctg cgg cgg gat ttg gtc gga Cys Leu Ser Phe Ala Met Ala Lys Ile Leu Arg Arg Asp Leu Val Gly 320 325 330 335	1007
gat gaa ggg gtg ttt cgg gca gct gag gcg atc gcg gaa ata gag Asp Glu Gly Val Phe Arg Ala Ala Glu Ala Ile Ala Glu Ile Glu 340 345 350	1055
aag agg acg agc gac aag aag att cta gaa act gtg gag aac tgg ccg Lys Arg Thr Ser Asp Lys Lys Ile Leu Glu Thr Val Glu Asn Trp Pro 355 360 365	1103
tct gag att cgc gaa gcc ttg caa aac tgt tat ttc tcg gtg gcg gga Ser Glu Ile Arg Glu Ala Leu Gln Asn Cys Tyr Phe Ser Val Ala Gly 370 375 380	1151
tcg agc agg ctt gat ctt tac ggc gcg gat ttt gga tgg ggt aag gcg Ser Ser Arg Leu Asp Leu Tyr Gly Ala Asp Phe Gly Trp Gly Lys Ala 385 390 395	1199
tgf aag caa gag ata ctg tcg att gat gga gag aag ttt acg atg tcg Val Lys Gln Glu Ile Leu Ser Ile Asp Gly Glu Lys Phe Thr Met Ser 400 405 410 415	1247
ttg tgt aaa ccg agg gat gct gcc gga gga ttg gag gtt gga ttg tct Leu Cys Lys Pro Arg Asp Ala Ala Gly Gly Leu Glu Val Gly Leu Ser 420 425 430	1295
ttg cca aag gag gaa ttg caa gct ttt gat gat tat ttt gcg gag gga Leu Pro Lys Glu Glu Leu Gln Ala Phe Asp Asp Tyr Phe Ala Glu Gly 435 440 445	1343
ata aag ggt tgattaatca tttaatcatg tattatgaag ttggatgaaa Ile Lys Gly 450	1392
tcctctgttt catctctatt gtttaaacaa taattttttt ccattgaact ttttgagtc 1452	
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<212> PRT  
<213> Artificial Sequence

<220>  
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Arg Phe Leu Gly Ile Thr Gly Ser Pro Lys  
1 5 10

<210> 8  
<211> 8  
<212> PRT  
<213> Artificial Sequence

<220>  
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1 5

<210> 9  
<211> 10  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic peptide

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1 5 10

<210> 10  
<211> 8  
<212> PRT  
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<220>  
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1 5

<210> 11  
<211> 14  
<212> PRT  
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<220>  
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<400> 11  
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<210> 12  
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<220>  
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1 5 10

<210> 13  
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<212> PRT  
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1 5

<210> 14  
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<212> PRT  
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1 5

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1 5

<210> 16  
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<222> (18)  
<223> a, c, g, t, unknown, or other

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23

<210> 17  
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23

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26

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<210> 19
<211> 17
<212> DNA
<213> Artificial Sequence

<220>
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      oligonucleotide

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<400> 19
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17

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<210> 20
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<213> Artificial Sequence

<220>
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      oligonucleotide

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<400> 20
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16

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<210> 21
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<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic
      peptide

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<400> 21
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    1           5

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<210> 22
<211> 17
<212> DNA
<213> Artificial Sequence

<220>
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      oligonucleotide

<220>
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<222> (9)
<223> a, c, g, t, unknown, or other

<220>
<221> modified_base
<222> (15)
<223> a, c, g, t, unknown, or other

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<210> 23 <211> 21 <212> DNA <213> Artificial Sequence	
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<210> 27
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<212> DNA
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<220>
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<220>
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<222> (12)..(53)

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      1           5           10

aga                                53
Arg

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<210> 28
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<212> DNA
<213> Artificial Sequence

<220>
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      oligonucleotide

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<210> 29
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<212> DNA
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<220>
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<222> (12)..(44)

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      Met Glu Gln Ile Gln Met Val Asn Ile Leu Glu
      1           5           10

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<210> 30  
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 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
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 oligonucleotide

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21

<210> 31  
 <211> 35  
 <212> DNA  
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 <223> Description of Artificial Sequence: Synthetic  
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 Thr Thr Leu Leu Glu Ser  
 1 5

35

<210> 32  
 <211> 469  
 <212> PRT  
 <213> Gentiana triflora

<400> 32  
 Met Glu Gln Ile Gln Met Val Lys Val Leu Glu Lys Cys Gln Val Thr  
 1 5 10 15

Pro Pro Ser Asp Thr Thr Asp Val Glu Leu Ser Leu Pro Val Thr Phe  
 20 25 30

Phe Asp Ile Pro Trp Leu His Leu Asn Lys Met Gln Ser Leu Leu Phe  
 35 40 45

Tyr Asp Phe Pro Tyr Pro Arg Thr His Phe Leu Asp Thr Val Ile Pro  
 50 55 60

Asn Leu Lys Ala Ser Leu Ser Leu Thr Leu Lys His Tyr Val Pro Leu  
 65 70 75 80

Ser Gly Asn Leu Leu Met Pro Ile Lys Ser Gly Glu Met Pro Lys Phe  
 85 90 95

Gln Tyr Ser Arg Asp Glu Gly Asp Ser Ile Thr Leu Ile Val Ala Glu  
     100                       105                       110  
  
 Ser Asp Gln Asp Phe Asp Tyr Leu Lys Gly His Gln Leu Val Asp Ser  
     115                       120                       125  
  
 Asn Asp Leu His Gly Leu Phe Tyr Val Met Pro Arg Val Ile Arg Thr  
     130                       135                       140  
  
 Met Gln Asp Tyr Lys Val Ile Pro Leu Val Ala Val Gln Val Thr Val  
     145                       150                       155                       160  
  
 Phe Pro Asn Arg Gly Ile Ala Val Ala Leu Thr Ala His His Ser Ile  
     165                       170                       175  
  
 Ala Asp Ala Lys Ser Phe Val Met Phe Ile Asn Ala Trp Ala Tyr Ile  
     180                       185                       190  
  
 Asn Lys Phe Gly Lys Asp Ala Asp Leu Leu Ser Ala Asn Leu Leu Pro  
     195                       200                       205  
  
 Ser Phe Asp Arg Ser Ile Ile Lys Asp Leu Tyr Gly Leu Glu Glu Thr  
     210                       215                       220  
  
 Phe Trp Asn Glu Met Gln Asp Val Leu Glu Met Phe Ser Arg Phe Gly  
     225                       230                       235                       240  
  
 Ser Lys Pro Pro Arg Phe Asn Lys Val Arg Ala Thr Tyr Val Leu Ser  
     245                       250                       255  
  
 Leu Ala Glu Ile Gln Lys Leu Lys Asn Lys Val Leu Asn Leu Arg Gly  
     260                       265                       270  
  
 Ser Glu Pro Thr Ile Arg Val Thr Thr Phe Thr Met Thr Cys Gly Tyr  
     275                       280                       285  
  
 Val Trp Thr Cys Met Val Lys Ser Lys Asp Asp Val Val Ser Glu Glu  
     290                       295                       300  
  
 Ser Ser Asn Asp Glu Asn Glu Leu Glu Tyr Phe Ser Phe Thr Ala Asp  
     305                       310                       315                       320  
  
 Cys Arg Gly Leu Leu Thr Pro Pro Cys Pro Pro Asn Tyr Phe Gly Asn  
     325                       330                       335  
  
 Cys Leu Ala Ser Cys Val Ala Lys Ala Thr His Lys Glu Leu Val Gly  
     340                       345                       350  
  
 Asp Lys Gly Leu Leu Val Ala Val Ala Ala Ile Gly Glu Ala Ile Glu  
     355                       360                       365  
  
 Lys Arg Leu His Asn Glu Lys Gly Val Leu Ala Asp Ala Lys Thr Trp  
     370                       375                       380  
  
 Leu Ser Glu Ser Asn Gly Ile Pro Ser Lys Arg Phe Leu Gly Ile Thr  
     385                       390                       395                       400

Gly Ser Pro Lys Phe Asp Ser Tyr Gly Val Asp Phe Gly Trp Gly Lys  
 405 410 415

Pro Ala Lys Phe Asp Ile Thr Ser Val Asp Tyr Ala Glu Leu Ile Tyr  
 420 425 430

Val Ile Gln Ser Arg Asp Phe Glu Lys Gly Val Glu Ile Gly Val Ser  
 435 440 445

Leu Pro Lys Ile His Met Asp Ala Phe Ala Lys Ile Phe Glu Glu Gly  
 450 455 460

Phe Cys Ser Leu Ser  
 465

<210> 33  
 <211> 479  
 <212> PRT  
 <213> Gentiana triflora

<400> 33  
 Met Ala Gly Asn Ser Glu Asp Ile Lys Val Leu Glu Lys Cys Arg Val  
 1 5 10 15

Ala Pro Pro Pro Asp Ala Val Ala Glu Phe Thr Val Pro Leu Ser Phe  
 20 25 30

Phe Asp Met Arg Trp Leu Ile Ser Asp Ala Glu His His Leu His Phe  
 35 40 45

Tyr Arg Phe Arg His Pro Cys Pro Asn Ser Lys Phe Ile Ile Ser Ser  
 50 55 60

Ile Lys Ser Ser Leu Ser Leu Val Leu Lys His Phe Leu Pro Leu Ala  
 65 70 75 80

Gly Asn Leu Ile Trp Pro Val Asp Ser Ser Asp Arg Met Pro Glu Leu  
 85 90 95

Arg Tyr Lys Lys Gly Asp Ser Val Ser Leu Thr Ile Ala Glu Ser Ser  
 100 105 110

Met Asp Phe Asp Tyr Leu Ala Gly Asp His Gln Arg Asp Ser Tyr Lys  
 115 120 125

Phe Asn Asp Leu Ile Pro Gln Leu Pro Glu Pro Ile Val Thr Ser Gly  
 130 135 140

Asp Glu Val Leu Pro Leu Phe Ala Leu Gln Val Thr Val Phe Ser Asn  
 145 150 155 160

Thr Gly Ile Cys Ile Gly Arg Asn Leu His Gln Val Leu Gly Asp Ala  
 165 170 175

Ser Ser Phe Leu His Phe Asn Lys Leu Trp Val Leu Val Asp Lys Ser  
 180 185 190

Asn Gly Asp Ser Leu Lys Phe Leu Pro Leu Ser Ser Leu Pro Met Tyr  
 195 200 205

Asp Arg Ser Val Val Gln Asp Pro Phe His Ile Arg Arg Lys Ile Tyr  
 210 215 220

Asn Glu Arg Lys Leu Leu Lys Ser Gln Gly Thr Pro Thr Val Leu Asn  
 225 230 235 240

Pro Ala Ile Ser Lys Asp Glu Val Arg Ala Thr Phe Ile Leu His Pro  
 245 250 255

Ile Asp Ile Met Lys Leu Lys Lys Phe Ile Ser Ser Lys Asn Arg Asn  
 260 265 270

Leu Thr Gly Ser Ser Asn Tyr Asn Leu Ser Thr Phe Thr Val Thr Ser  
 275 280 285

Ala Leu Ile Trp Thr Cys Leu Ser Lys Ser Leu Asp Thr Val Val Arg  
 290 295 300

Glu Lys Val Glu Glu Asp Lys His Ala Ala Asn Leu Cys Ala Phe Ile  
 305 310 315 320

Asn Cys Arg Gln Arg Phe Ala Pro Pro Ile Pro Gln Asn Tyr Phe Gly  
 325 330 335

Asn Cys Ile Val Pro Cys Met Val Gly Ser Thr His Glu Gln Leu Val  
 340 345 350

Gly Asn Glu Gly Leu Ser Val Ala Ala Thr Ala Ile Gly Asp Ala Ile  
 355 360 365

His Lys Arg Leu His Asp Tyr Glu Gly Ile Leu Arg Gly Asp Trp Ile  
 370 375 380

Ser Pro Pro Arg Ser Thr Ser Ala Ala Pro Arg Ser Thr Leu Ile Tyr  
 385 390 395 400

Val Val Gly Ser Ala Gln Arg Asn Val His Asp Phe Asp Ala Asp Phe  
 405 410 415

Gly Trp Gly Lys Leu Glu Lys His Glu Ser Val Ser Thr Asn Pro Ser  
 420 425 430

Ala Thr Leu Ile Leu Ile Ser Arg Ser Arg Arg Phe Lys Gly Ala Leu  
 435 440 445

Glu Leu Gly Ile Ser Leu Pro Lys Asn Arg Met Asp Ala Phe Ala Thr  
 450 455 460

Ile Phe Thr Asn Phe Ile Asn Ser Leu His Val Arg Ser Pro Leu  
 465 470 475

&lt;210&gt; 34

&lt;211&gt; 448

&lt;212&gt; PRT

&lt;213&gt; Petunia hybrida

&lt;400&gt; 34

Met Ala Gly Glu Val Ala Lys Gln Glu Val Thr Lys Val Lys Val Leu				
1	5	10	15	

Lys Lys Thr Asn Val Lys Pro His Lys Pro Leu Gly Lys Lys Glu Cys				
20	25	30		

Gln Leu Val Thr Phe Asp Leu Pro Tyr Leu Ala Phe Tyr Tyr Asn Gln				
35	40	45		

Lys Phe Leu Ile Tyr Lys Gly Ala Glu Asn Phe Asp Glu Thr Val Glu				
50	55	60		

Lys Ile Lys Asp Gly Leu Ala Leu Val Leu Val Asp Phe Tyr Gln Leu				
65	70	75	80	

Ala Gly Lys Leu Gly Lys Asp Glu Glu Gly Val Phe Arg Val Glu Tyr				
85	90	95		

Asp Asp Asp Met Asp Gly Val Glu Val Thr Val Ala Val Ala Glu Glu				
100	105	110		

Ile Glu Val Ala Asp Leu Thr Asp Glu Glu Gly Thr Thr Lys Leu Gln				
115	120	125		

Asp Leu Ile Pro Cys Asn Lys Ile Leu Asn Leu Glu Gly Leu His Arg				
130	135	140		

Pro Leu Leu Ala Val Gln Leu Thr Lys Leu Lys Asp Gly Leu Thr Met				
145	150	155	160	

Gly Leu Ala Phe Asn His Ala Val Leu Asp Gly Thr Ser Thr Trp His				
165	170	175		

Phe Met Thr Ser Trp Ser Glu Leu Cys Cys Gly Ser Thr Ser Ile Ser				
180	185	190		

Val Pro Pro Phe Leu Glu Arg Thr Lys Ala Arg Asn Thr Arg Val Lys				
195	200	205		

Leu Asn Leu Ser Gln Pro Ser Asp Ala Pro Glu His Ala Lys Ser Ala				
210	215	220		

Thr Asn Gly Asp Val Pro Ala Asn Val Asp Pro Pro Leu Arg Glu Arg				
225	230	235	240	

Val Phe Lys Phe Ser Glu Leu Ala Ile Asp Lys Ile Lys Ser Thr Val				
245	250	255		

Asn Ala Asn Ser Gly Glu Thr Pro Phe Ser Thr Phe Gln Ser Leu Ser				
260	265	270		

Ala His Val Trp Leu Ala Val Thr Arg Ala Arg Gln Leu Lys Pro Glu  
 275                    280                    285  
  
 Asp Tyr Thr Val Tyr Thr Val Phe Ala Asp Cys Arg Lys Arg Val Asp  
 290                    295                    300  
  
 Pro Pro Met Pro Glu Ser Tyr Phe Gly Asn Leu Ile Gln Ala Ile Phe  
 305                    310                    315                    320  
  
 Thr Val Thr Ala Ala Gly Leu Leu Leu Ala Ser Pro Ile Glu Phe Ala  
 325                    330                    335  
  
 Gly Gly Met Ile Gln Gln Ala Ile Val Lys His Asp Ala Lys Ala Ile  
 340                    345                    350  
  
 Asp Glu Arg Asn Lys Glu Trp Glu Ser Asn Pro Lys Ile Phe Gln Tyr  
 355                    360                    365  
  
 Lys Asp Ala Gly Val Asn Cys Val Ala Val Gly Ser Ser Pro Arg Phe  
 370                    375                    380  
  
 Lys Val Tyr Asp Val Asp Phe Gly Trp Gly Lys Pro Glu Ser Val Arg  
 385                    390                    395                    400  
  
 Ser Gly Ser Asn Asn Arg Phe Asp Gly Met Val Tyr Leu Tyr Gln Gly  
 405                    410                    415  
  
 Lys Asn Gly Gly Arg Ser Ile Asp Val Glu Ile Ser Leu Glu Ala Asn  
 420                    425                    430  
  
 Ala Met Glu Arg Leu Glu Lys Asp Lys Glu Phe Leu Met Glu Thr Ala  
 435                    440                    445

<210> 35  
 <211> 446  
 <212> PRT  
 <213> Perilla ocimoides

<400> 35  
 Val Ile Glu Thr Cys Arg Val Gly Pro Pro Pro Asp Ser Val Ala Glu  
 1                    5                    10                    15  
  
 Gln Ser Val Pro Leu Thr Phe Phe Asp Met Thr Trp Leu His Phe His  
 20                    25                    30  
  
 Pro Met Leu Gln Leu Leu Phe Tyr Glu Phe Pro Cys Ser Lys Gln His  
 35                    40                    45  
  
 Phe Ser Glu Ser Ile Val Pro Lys Leu Lys Gln Ser Leu Ser Lys Thr  
 50                    55                    60  
  
 Leu Ile His Phe Phe Pro Leu Ser Cys Asn Leu Ile Tyr Pro Ser Ser  
 65                    70                    75                    80  
  
 Pro Glu Lys Met Pro Glu Phe Arg Tyr Leu Ser Gly Asp Ser Val Ser  
 85                    90                    95

Phe Thr Ile Ala Glu Ser Ser Asp Asp Phe Asp Asp Leu Val Gly Asn  
 100 105 110

Arg Pro Glu Ser Pro Val Arg Leu Tyr Asn Phe Val Pro Lys Leu Pro  
 115 120 125

Pro Ile Val Glu Glu Ser Asp Arg Lys Leu Phe Gln Val Phe Ala Val  
 130 135 140

Gln Val Thr Leu Phe Pro Gly Arg Gly Val Gly Ile Gly Ile Ala Thr  
 145 150 155 160

His His Thr Val Ser Asp Ala Pro Ser Phe Leu Ala Phe Ile Thr Ala  
 165 170 175

Trp Ser Ser Met Ser Lys His Ile Glu Asn Glu Asp Glu Asp Glu Glu  
 180 185 190

Phe Lys Ser Leu Pro Val Phe Asp Arg Ser Val Ile Lys Tyr Pro Thr  
 195 200 205

Lys Phe Asp Ser Ile Tyr Trp Arg Asn Ala Leu Lys Phe Pro Leu Gln  
 210 215 220

Ser Arg His Pro Ser Leu Pro Thr Asp Arg Ile Arg Thr Thr Phe Val  
 225 230 235 240

Phe Thr Gln Ser Lys Ile Lys Lys Leu Lys Gly Trp Ile Gln Ser Arg  
 245 250 255

Val Pro Ser Leu Val His Leu Ser Ser Phe Val Ala Ile Ala Ala Tyr  
 260 265 270

Met Trp Ala Gly Ile Thr Lys Ser Phe Thr Ala Asp Glu Asp Gln Asp  
 275 280 285

Asn Glu Asp Ala Phe Phe Leu Ile Pro Val Asp Leu Arg Pro Arg Leu  
 290 295 300

Asp Pro Pro Val Pro Glu Asn Tyr Phe Gly Asn Cys Leu Ser Tyr Ala  
 305 310 315 320

Leu Pro Arg Met Arg Arg Glu Leu Val Gly Glu Lys Gly Val Phe  
 325 330 335

Leu Ala Ala Glu Val Ile Ala Ala Glu Ile Lys Lys Arg Ile Asn Asp  
 340 345 350

Lys Arg Ile Leu Glu Thr Val Glu Lys Trp Ser Pro Glu Ile Arg Lys  
 355 360 365

Ala Leu Gln Lys Ser Tyr Phe Ser Val Ala Gly Ser Ser Lys Leu Asp  
 370 375 380

Leu Tyr Gly Ala Asp Phe Gly Trp Gly Lys Ala Arg Lys Gln Glu Ile  
 385 390 395 400

Leu Ser Ile Asp Gly Glu Lys Tyr Ala Met Thr Leu Cys Lys Ala Arg  
                   405                  410                  415  
  
 Asp Phe Glu Gly Gly Leu Glu Val Cys Leu Ser Leu Pro Lys Asp Lys  
                   420                  425                  430  
  
 Met Asp Ala Phe Ala Ala Tyr Phe Ser Leu Gly Ile Asn Gly  
                   435                  440                  445  
  
  
 <210> 36  
 <211> 454  
 <212> PRT  
 <213> Senecio cruentus  
  
 <400> 36  
 Asn Ile Leu Glu His Ala Arg Ile Ser Ala Pro Ser Gly Thr Ile Gly  
       1                  5                  10                  15  
  
 His Arg Ser Leu Ser Leu Thr Phe Phe Asp Ile Thr Trp Leu Leu Phe  
       20                  25                  30  
  
 Pro Pro Val His His Leu Phe Phe Tyr Asp Phe Pro His Ser Lys Ser  
       35                  40                  45  
  
 His Phe Met Asp Thr Ile Val Pro Arg Leu Lys Gln Ser Leu Ser Val  
       50                  55                  60  
  
 Thr Leu Gln His Phe Phe Pro Phe Ala Ser Asn Leu Ile Val Phe Pro  
       65                  70                  75                  80  
  
 Asn Thr Asp Gly Ser Gly Phe Asn Lys Lys Pro Glu Ile Lys His Val  
       85                  90                  95  
  
 Glu Gly Asp Ser Val Val Val Thr Phe Ala Glu Cys Cys Leu Asp Phe  
       100                105                  110  
  
 Asn Asn Leu Thr Gly Asn His Pro Arg Lys Cys Glu Asn Phe Tyr Pro  
       115                120                  125  
  
 Leu Val Pro Ser Leu Gly Asn Ala Ile Lys Leu Cys Asp Cys Val Thr  
       130                135                  140  
  
 Val Pro Leu Phe Ser Leu Gln Val Thr Phe Phe Pro Gly Ser Gly Ile  
       145                150                  155                  160  
  
 Ser Leu Gly Met Thr Asn His His Ser Leu Gly Asp Ala Ser Thr Arg  
       165                170                  175  
  
 Phe Asn Phe Leu Lys Gly Trp Thr Ser Ile Ile Gln Ser Gly Val Asp  
       180                185                  190  
  
 Arg Ser Phe Leu Thr Lys Gly Ser Pro Pro Val Phe Asp Arg Leu Ile  
       195                200                  205  
  
 Asn Ile Pro His Leu Asp Glu Asn Lys Leu Arg His Thr Arg Leu Glu  
       210                215                  220

Ser Phe Tyr Lys Pro Ser Ser Leu Val Gly Pro Thr Asp Lys Val Arg  
 225                    230                    235                    240  
  
 Ser Thr Phe Val Leu Thr Arg Thr Asn Ile Asn Leu Leu Lys Lys Lys  
 245                    250                    255  
  
 Val Leu Thr Gln Val Pro Asn Leu Glu Tyr Met Ser Ser Phe Thr Val  
 260                    265                    270  
  
 Thr Cys Gly Tyr Ile Trp Ser Cys Ile Ala Lys Ser Leu Val Lys Ile  
 275                    280                    285  
  
 Gly Glu Arg Lys Gly Glu Asp Glu Leu Glu Gln Phe Ile Ile Thr Ile  
 290                    295                    300  
  
 Asp Cys Arg Ser Arg Leu Asp Pro Pro Ile Pro Thr Ala Tyr Phe Gly  
 305                    310                    315                    320  
  
 Asn Cys Gly Ala Pro Cys Val Pro Thr Leu Lys Asn Val Val Leu Thr  
 325                    330                    335  
  
 Thr Glu Asn Gly Tyr Ala Leu Gly Ala Lys Val Ile Gly Glu Ser Ile  
 340                    345                    350  
  
 Cys Lys Met Ile Tyr Asn Lys Asp Gly Ile Leu Lys Asp Ala Ala Arg  
 355                    360                    365  
  
 Trp His Glu Pro Phe Met Ile Pro Ala Arg Lys Ile Gly Val Ala Gly  
 370                    375                    380  
  
 Thr Pro Lys Leu Asn Leu Tyr Asp Phe Asp Phe Gly Trp Gly Lys Arg  
 385                    390                    395                    400  
  
 Ile Lys Tyr Glu Thr Val Ser Ile Asp Tyr Asn Thr Ser Ile Ser Ile  
 405                    410                    415  
  
 Asn Ala Ser Lys Thr Ser Ala Gln Asp Leu Glu Ile Gly Leu Ser Leu  
 420                    425                    430  
  
 Pro Ser Met Gln Met Glu Ala Phe Ser Ser Ile Phe Asp Glu Gly Leu  
 435                    440                    445  
  
 Glu Ser Gln Val Ser Leu  
 450  
  
 <210> 37  
 <211> 450  
 <212> PRT  
 <213> Lavandula angustifolia  
  
 <400> 37  
 Thr Thr Leu Leu Glu Ser Ser Arg Val Ala Pro Pro Pro Gly Thr Val  
 1                    5                    10                    15  
  
 Ala Glu Gln Ser Leu Pro Leu Thr Phe Phe Asp Met Thr Trp Leu His  
 20                    25                    30

Phe His Pro Met Leu Gln Leu Leu Phe Tyr Glu Leu Pro Cys Ser Lys  
     35                  40                  45

Pro Ala Phe Leu Glu Thr Val Val Pro Lys Leu Lys Gln Ser Leu Ser  
     50                  55                  60

Leu Thr Leu Lys His Phe Phe Pro Leu Ser Cys Asn Leu Ile Tyr Pro  
     65                  70                  75                  80

Leu Ser Pro Glu Lys Met Pro Glu Phe Arg Tyr Gln Asn Gly Asp Ser  
     85                  90                  95

Val Ser Phe Thr Ile Met Glu Ser Val Gly Asp His Pro His Ser Ala  
     100                 105                 110

His Lys Tyr Tyr Cys Phe Ala Pro Ser Asp Asp Tyr Glu Asp Leu Gln  
     115                 120                 125

Leu Pro Pro Ile Val Glu Glu Ser Asp Arg Lys Leu Phe Gln Val Leu  
     130                 135                 140

Ala Val Gln Val Thr Leu Phe Pro Gly Arg Gly Val Cys Ile Gly Ile  
     145                 150                 155                 160

Thr Thr His His Thr Val Ser Asp Ala Pro Ser Phe Val Gly Phe Met  
     165                 170                 175

Lys Ser Trp Ala Ser Ile Thr Lys Phe Gly Gly Asp Asp Glu Phe Leu  
     180                 185                 190

Asp Gly Lys Gly Glu Cys Leu Pro Val Phe Asp Arg Ser Leu Val Asn  
     195                 200                 205

Tyr Pro Pro Lys Leu Asp Thr Tyr Leu Trp Asn Asn Ala Gln Lys Arg  
     210                 215                 220

Pro Leu Glu Ser Gln His Pro Ser Leu Pro Thr Asp Arg Ile Arg Ala  
     225                 230                 235                 240

Thr Tyr Leu Phe Thr Gln Ser Glu Ile Lys Lys Leu Lys Gly Leu Ile  
     245                 250                 255

Gln Arg Lys Ala Pro Asn Val Val Asn Leu Ser Ser Phe Val Ala Ile  
     260                 265                 270

Ala Ala Tyr Ile Trp Thr Gly Ile Ala Lys Ser Val Gly Asp Tyr Lys  
     275                 280                 285

Asp Val Asp Asp Asp Lys Arg Ala Phe Phe Leu Ile Pro Ile Asp Leu  
     290                 295                 300

Arg Pro Arg Leu Asp Pro Pro Ala Pro Gly Asn Tyr Phe Gly Asn Cys  
     305                 310                 315                 320

Leu Ser Phe Ala Met Ala Lys Ile Leu Arg Arg Asp Leu Val Gly Asp  
     325                 330                 335

Glu Gly Val Phe Arg Ala Ala Glu Ala Ile Ala Ala Glu Ile Glu Lys  
                  340                     345                     350

Arg Thr Ser Asp Lys Lys Ile Leu Glu Thr Val Glu Asn Trp Pro Ser  
                  355                     360                     365

Glu Ile Arg Glu Ala Leu Gln Asn Cys Tyr Phe Ser Val Ala Gly Ser  
                  370                     375                     380

Ser Arg Leu Asp Leu Tyr Gly Ala Asp Phe Gly Trp Gly Lys Ala Val  
                  385                     390                     395                     400

Lys Gln Glu Ile Leu Ser Ile Asp Gly Glu Lys Phe Thr Met Ser Leu  
                  405                     410                     415

Cys Lys Pro Arg Asp Ala Ala Gly Gly Leu Glu Val Gly Leu Ser Leu  
                  420                     425                     430

Pro Lys Glu Glu Leu Gln Ala Phe Asp Asp Tyr Phe Ala Glu Gly Ile  
                  435                     440                     445

Lys Gly  
         450

&lt;210&gt; 38

&lt;211&gt; 14

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Description of Artificial Sequence: Synthetic peptide

&lt;400&gt; 38

Met	Glu	Gln	Ile	Gln	Met	Val	Ala	Val	Ile	Glu	Thr	Cys	Arg
1					5								

10

&lt;210&gt; 39

&lt;211&gt; 11

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Description of Artificial Sequence: Synthetic peptide

&lt;400&gt; 39

Met	Glu	Gln	Ile	Gln	Met	Val	Asn	Ile	Leu	Glu
1					5					

10

&lt;210&gt; 40

&lt;211&gt; 6

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

<220>  
<223> Description of Artificial Sequence: Synthetic  
peptide

<400> 40  
Thr Thr Leu Leu Glu Ser  
1 5